Natural Computing Series

Nataša Jonoska Masahico Saito Editors

Discrete and Topological Models in Molecular Biology



<u>Discrete And Topological Models In Molecular Biology</u> <u>Natural Computing Series</u>

David Kirk

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Discrete and Topological Models in Molecular Biology Nataša Jonoska, Masahico Saito, 2013-12-23 Theoretical tools and insights from discrete mathematics theoretical computer science and topology now play essential roles in our understanding of vital biomolecular processes. The related methods are now employed in various fields of mathematical biology as instruments to zoom in on processes at a molecular level This book contains expository chapters on how contemporary models from discrete mathematics in domains such as algebra combinatorics and graph and knot theories can provide perspective on biomolecular problems ranging from data analysis molecular and gene arrangements and structures and knotted DNA embeddings via spatial graph models to the dynamics and kinetics of molecular interactions The contributing authors are among the leading scientists in this field and the book is a reference for researchers in mathematics and theoretical computer science who are engaged with modeling molecular and biological phenomena using discrete methods It may also serve as a guide and supplement for graduate courses in mathematical biology or bioinformatics introducing nontraditional aspects of mathematical biology **ReCombinatorics** Dan Gusfield, 2014-07-03 When a Philadelphia girl intercepts a message about an impending British attack against her father s regiment commanded by General Washington in White Marsh she travels alone by horseback to warn the Patriot army Integer Linear Programming in Computational and Systems Biology Dan Gusfield, 2019-06-13 This hands on tutorial text for non experts demonstrates biological applications of a versatile modeling and optimization technique Language, Life, Limits Arnold Beckmann, Ersébet Csuhaj-Varjú, Klaus Meer, 2014-06-05 This book constitutes the refereed proceedings of the 10th Conference on Computability in Europe CiE 2014 held in Budapest Hungary in June 2014 The 42 revised papers presented were carefully reviewed and selected from 78 submissions and included together with 15 invited papers in this proceedings. The conference had six special sessions computational linguistics bio inspired computation history and philosophy of computing computability theory online algorithms and complexity in automata theory Big Data and Differential Privacy Nii O. Attoh-Okine, 2017-05-12 A comprehensive introduction to the theory and practice of contemporary data science analysis for railway track engineering Featuring a practical introduction to state of the art data analysis for railway track engineering Big Data and Differential Privacy Analysis Strategies for Railway Track Engineering addresses common issues with the implementation of big data applications while exploring the limitations advantages and disadvantages of more conventional methods. In addition the book provides a unifying approach to analyzing large volumes of data in railway track engineering using an array of proven methods and software technologies Dr Attoh Okine considers some of today s most notable applications and implementations and highlights when a particular method or algorithm is most appropriate Throughout the book presents numerous real world examples to illustrate the latest railway engineering big data applications of predictive analytics such as the Union Pacific Railroad s use of big data to reduce train derailments increase the velocity of shipments and reduce emissions In

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as well as statistical methods for the solution of fundamental problems in the biosciences Describes how to find regularities among empirical data as well as conceptual models and theories

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